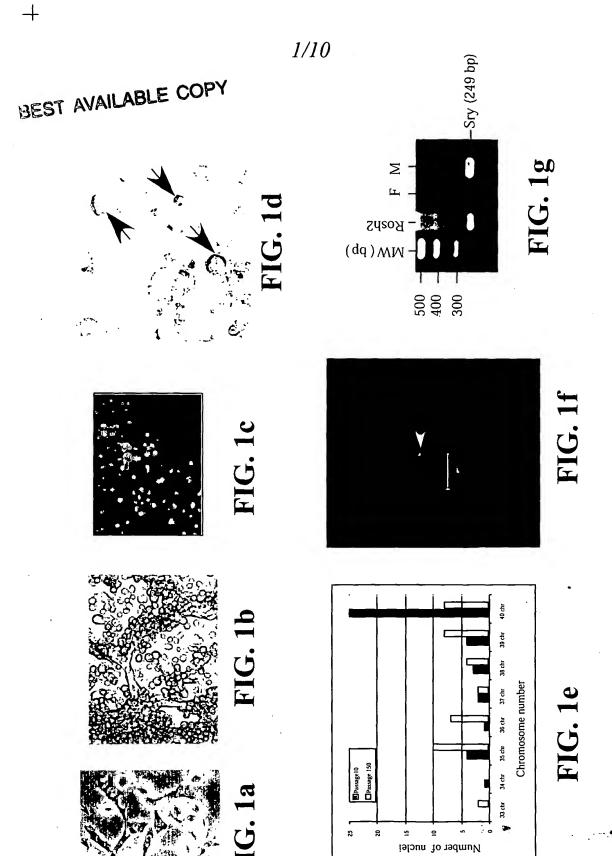
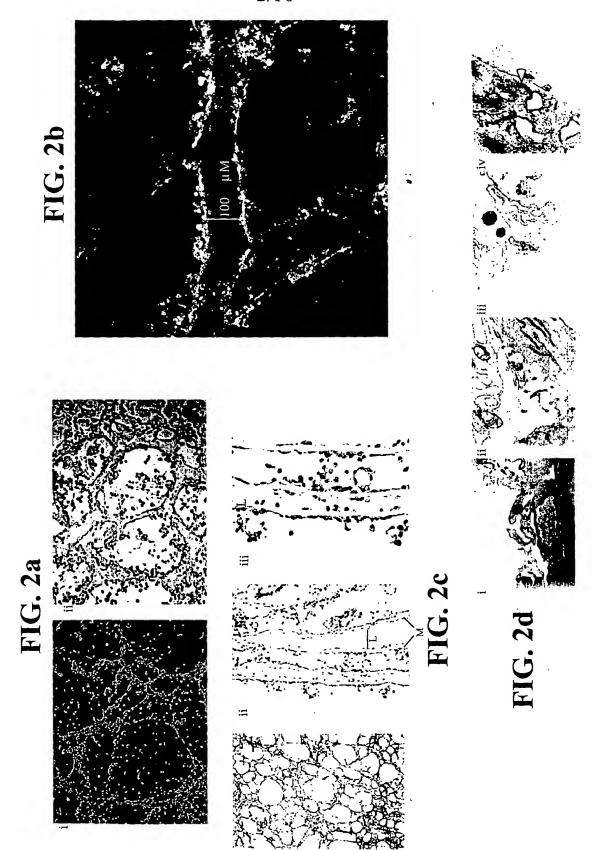
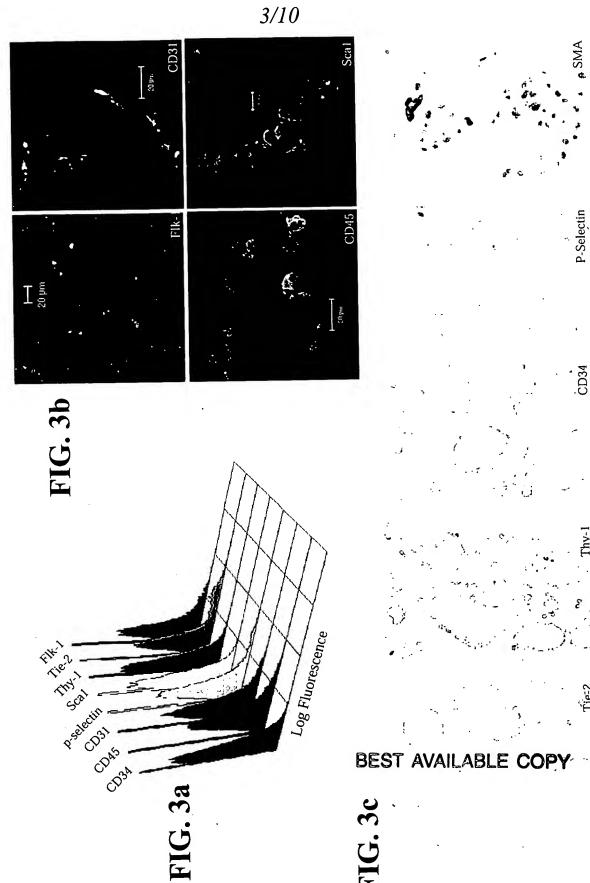
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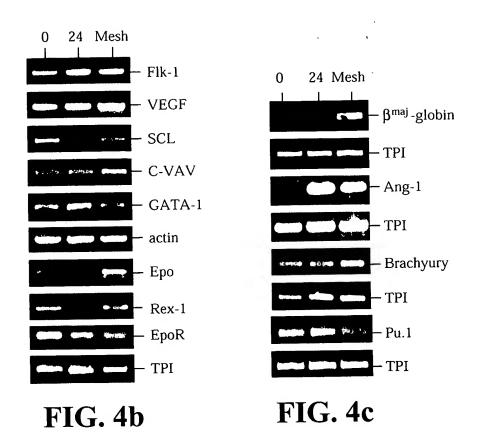
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Lower Primer CGTCGTATTCCTGTTTGCTG GCAACAACGAGGACATTAG GGAGGACTGGAGAAATCAG TGCTAGCCAATTCCTCCCAG GATGAGACCAGAATCTCTTTTC AGTCTCTGTAGGCCTCAGCT CACCCATCAAATCTCTTTTC AGTGTCTGTAGGCCTCAGCT CACCCATCAAAAGGACACAC CATCCAGTACGACACTGCTT GTTCTTTTTTTGTTTT GTTCTTTTTTTTTT	Lower Primer ACTTGCTGCTGCAACGGAGAC TGAAATCAGCACCGTGTAAG TGCTGCCTGTGAGTCATAAC GCAACAACGGAGGACATTAG	CCT CGC; CTG TCT TGC CGT AGT T	GTCCCCATGGAGTCAAAGAG CTCAAGGAGACCTTTGCTCA	CAGACCCCAGAAACAGACA CAGGCAAGAAGAGGGG
SEQ ID NO: 4 SEQ ID NO: 6 SEQ ID NO: 8 SEQ ID NO: 10 SEQ ID NO: 12 SEQ ID NO: 14 SEQ ID NO: 16 SEQ ID NO: 18 SEQ ID NO: 20 SEQ ID NO: 22 SEQ ID NO: 22	ID NO: ID NO: ID NO: ID NO:	SEQ ID NO: 34 SEQ ID NO: 36	SEQ ID NO: 38	SEQ ID NO: 42 SEQ ID NO: 44
Upper Primer CCGCCTCCTCCCTG GAAGCCAAGGACAC GTATAACGTGGAGGTCAAGC ATGTCACTGCTGGTGCTGGA CCTATGACCACCCACCGCATCG CGCTCTGTGGTTCTGCGTG ACT GAT ACC ACT AAG GT ACTGACCCTAAAGCAAGACG ATTGCACACGGGATTCTG CCCTGGCATGATCAAAGACTT TGAACTTTCTGCTCTTGGG		GAG ACA GGC AGC AAG AAA AAG GGA CAG CAT CTG GTG GGT GGA C	aaggaaagaaggaaag gtccccatggagtcaaaga	CTGACAGATGCTCTTTGGG TGACTGATGCTGAAGGCTG
SEQ ID NO: 3 SEQ ID NO: 5 SEQ ID NO: 7 SEQ ID NO: 11 SEQ ID NO: 13 SEQ ID NO: 15 SEQ ID NO: 17 SEQ ID NO: 17 SEQ ID NO: 21 SEQ ID NO: 23	ID NO: ID NO: ID NO: ID NO:	SEQ ID NO: 33	SEQ ID NO: 37	SEQ ID NO: 41
RT-PCR Size Actin 402 3rachyury 194 2-VAV 604 3po 409 3poR 702 3kr.1 500 3ATA-1 263 3cL 744 3PI 523 /EGF 406	Vene Size Vng-1 770 Srachyury 947	'U.1 648 164	H1 globin 792 265	maj globin 578 457

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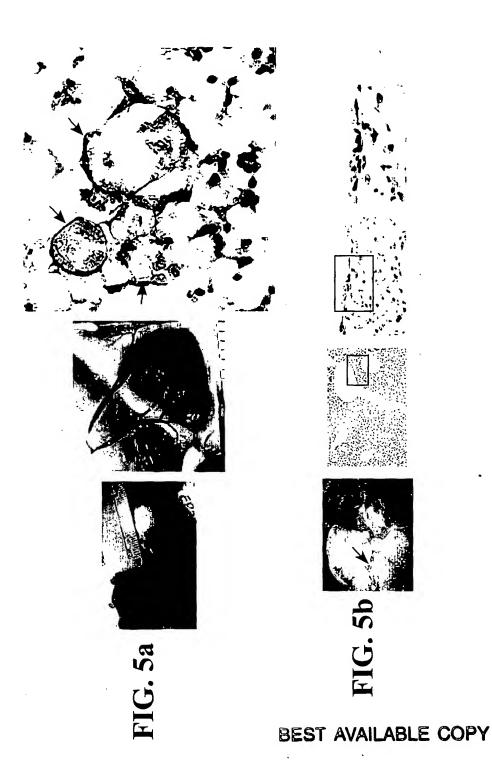
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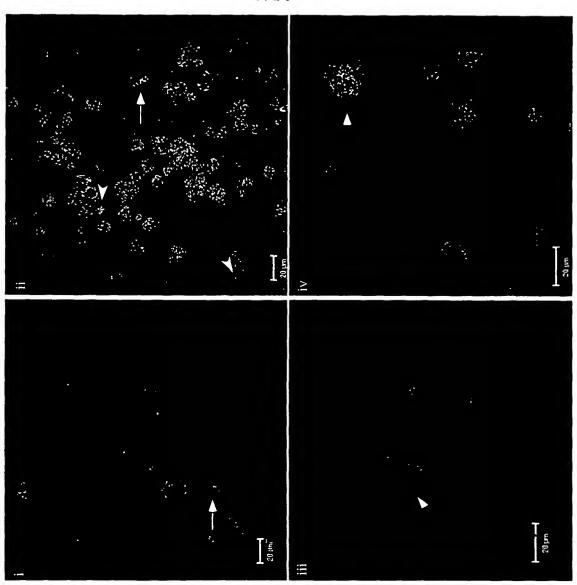
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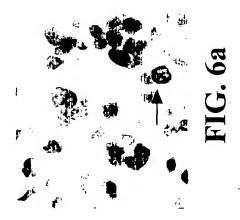
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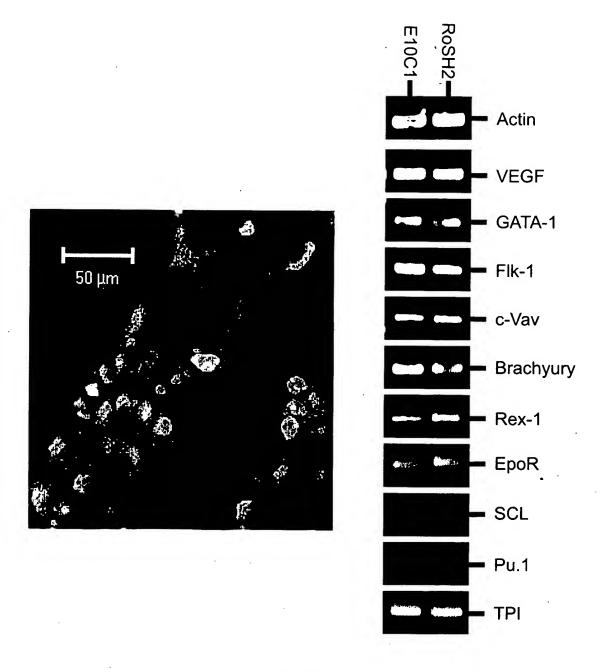


FIG. 7

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FIG. 8

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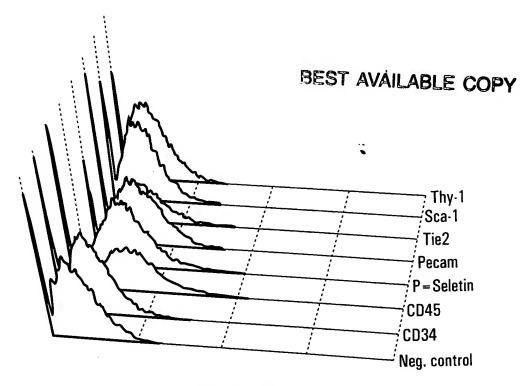


FIG. 9

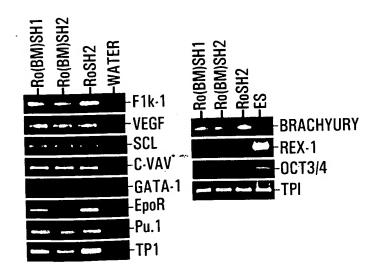


FIG. 10